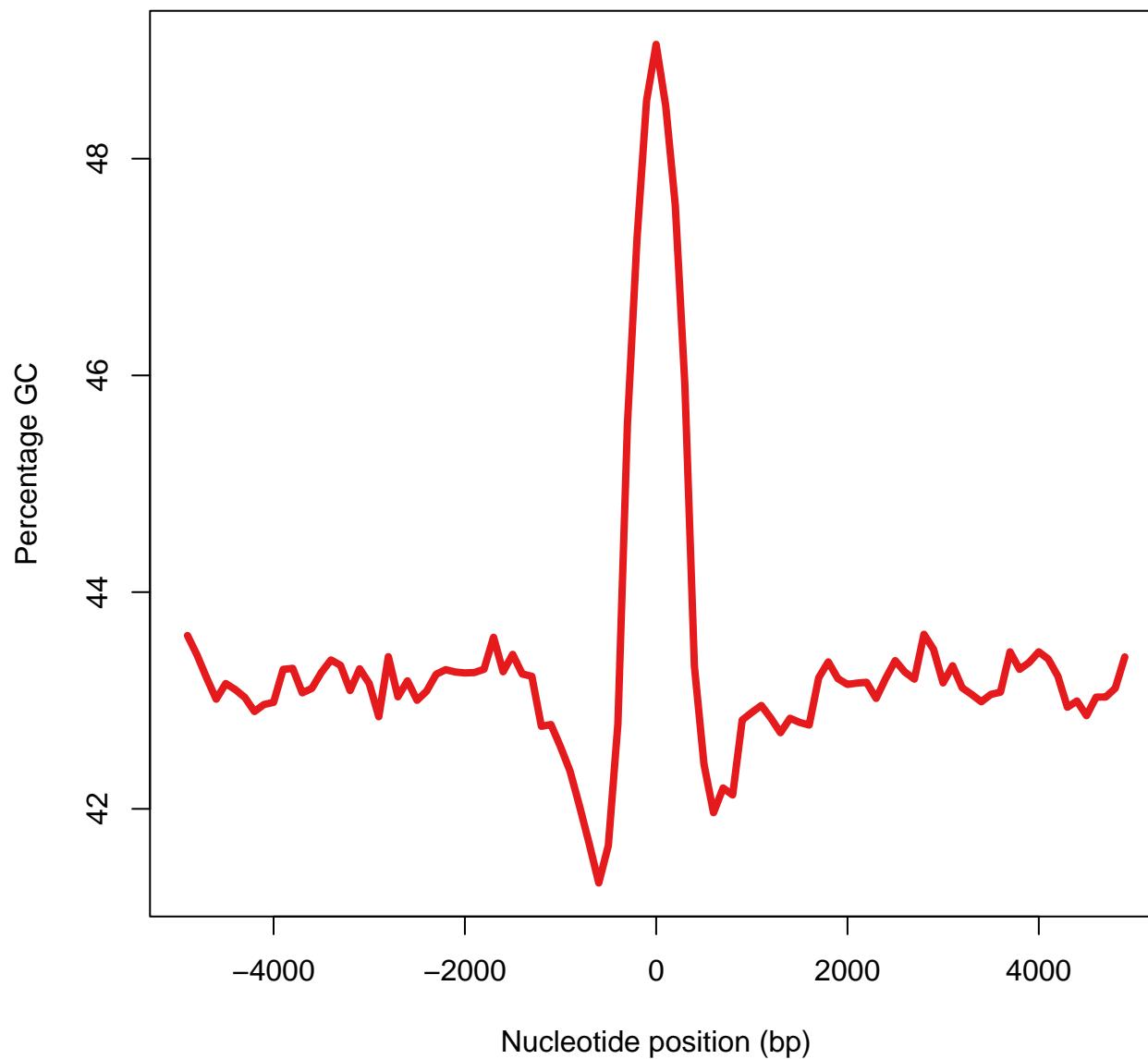
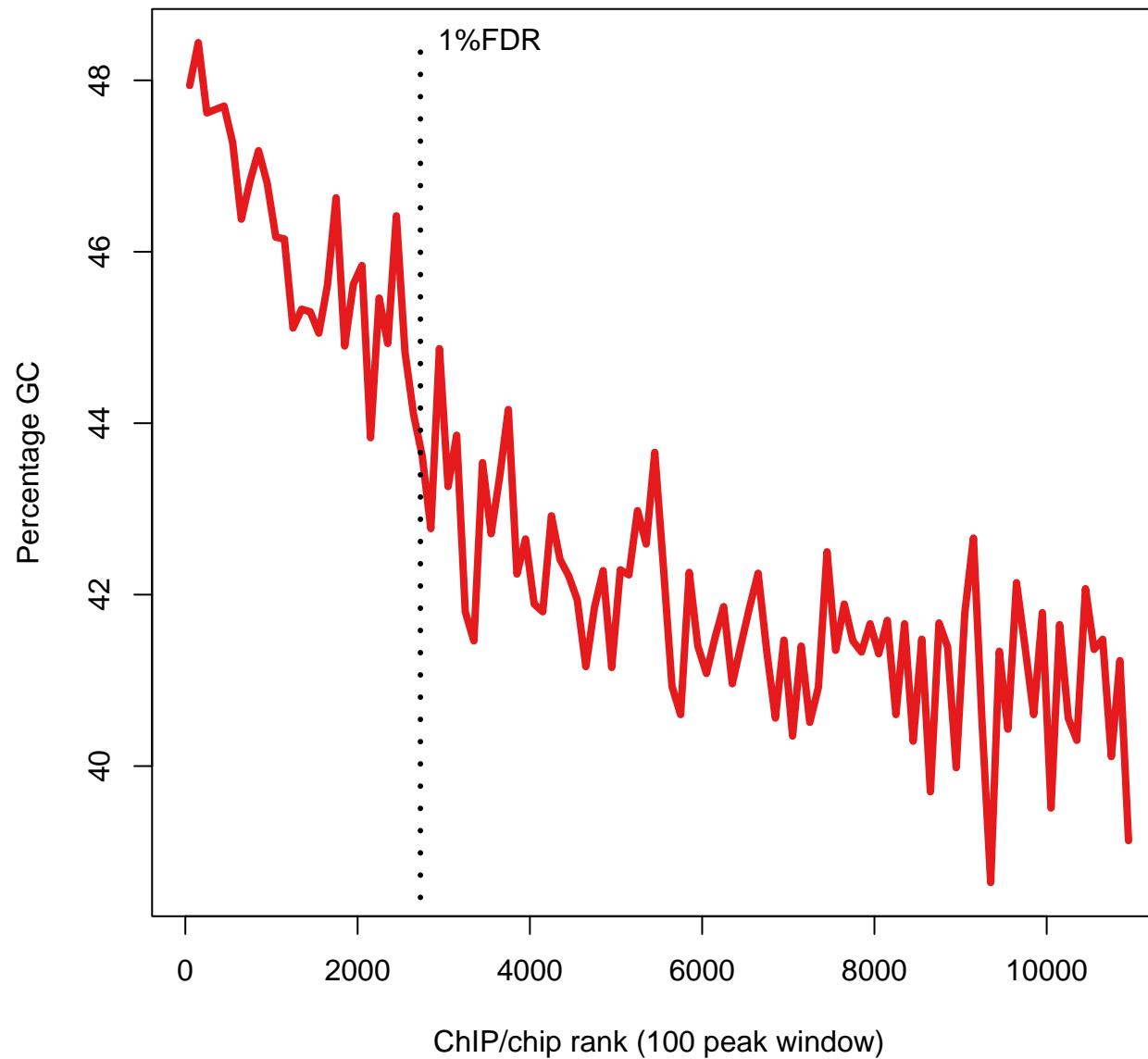


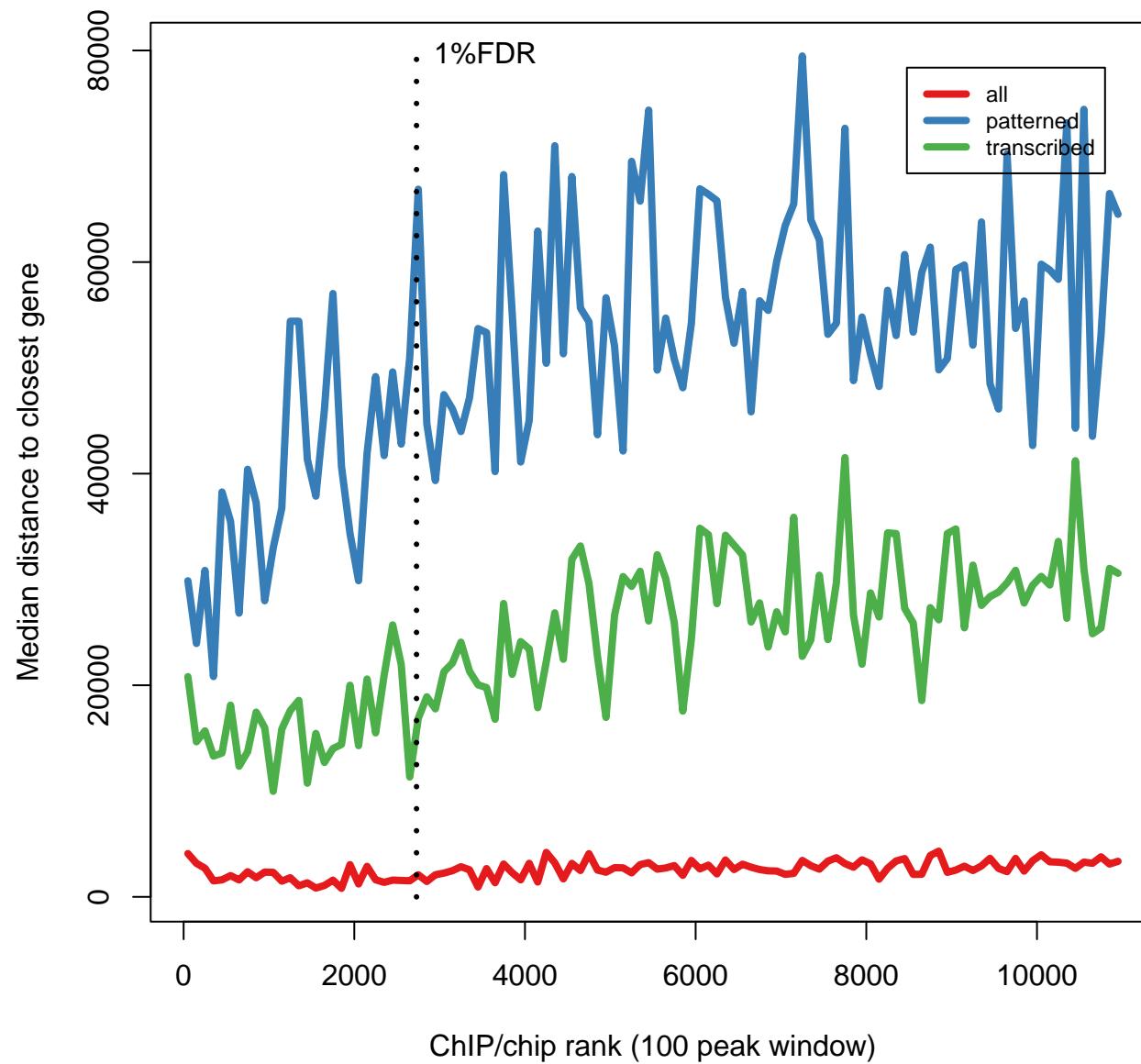
h_2 base composition across peaks



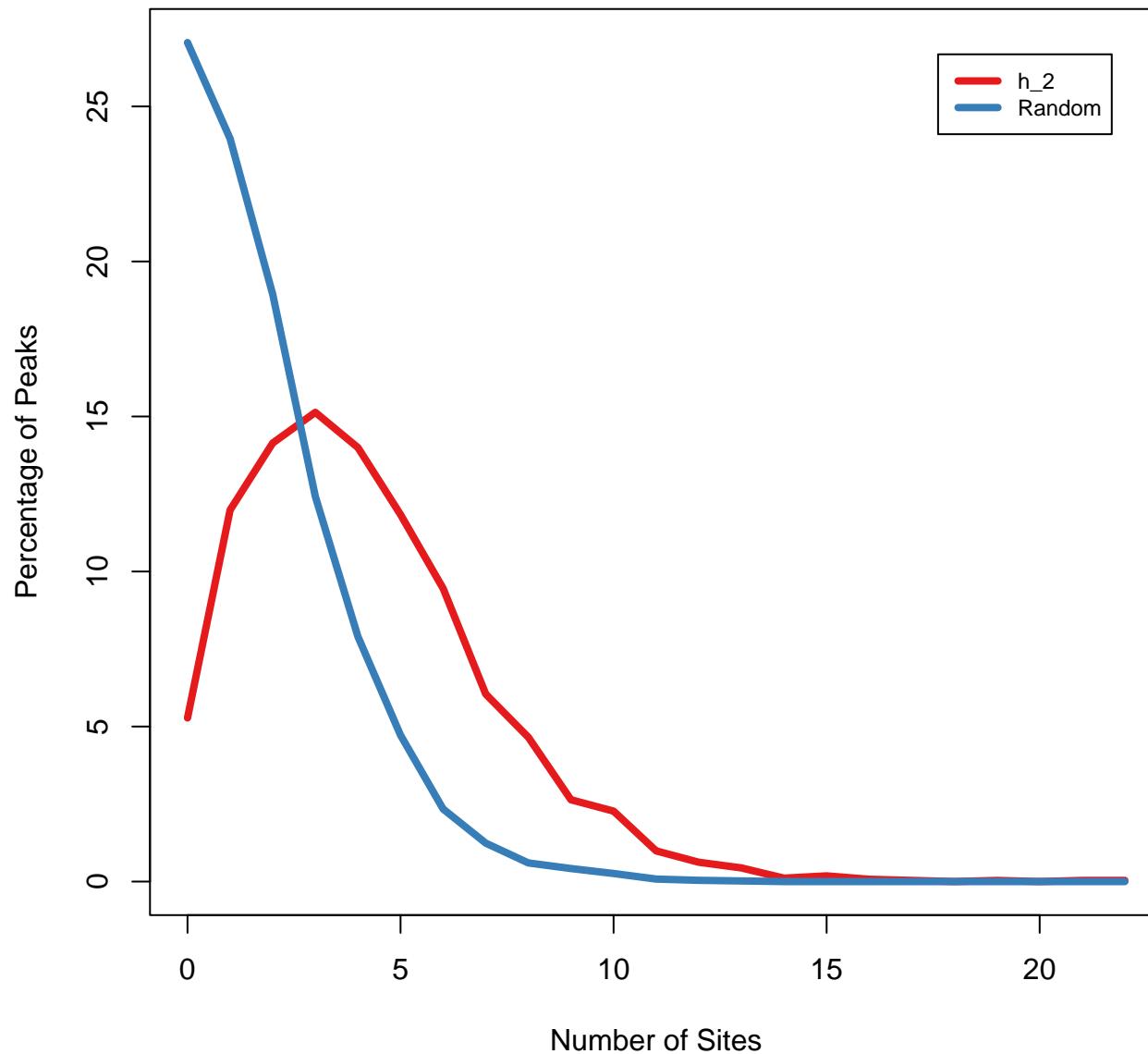
h_2 base composition down ranks



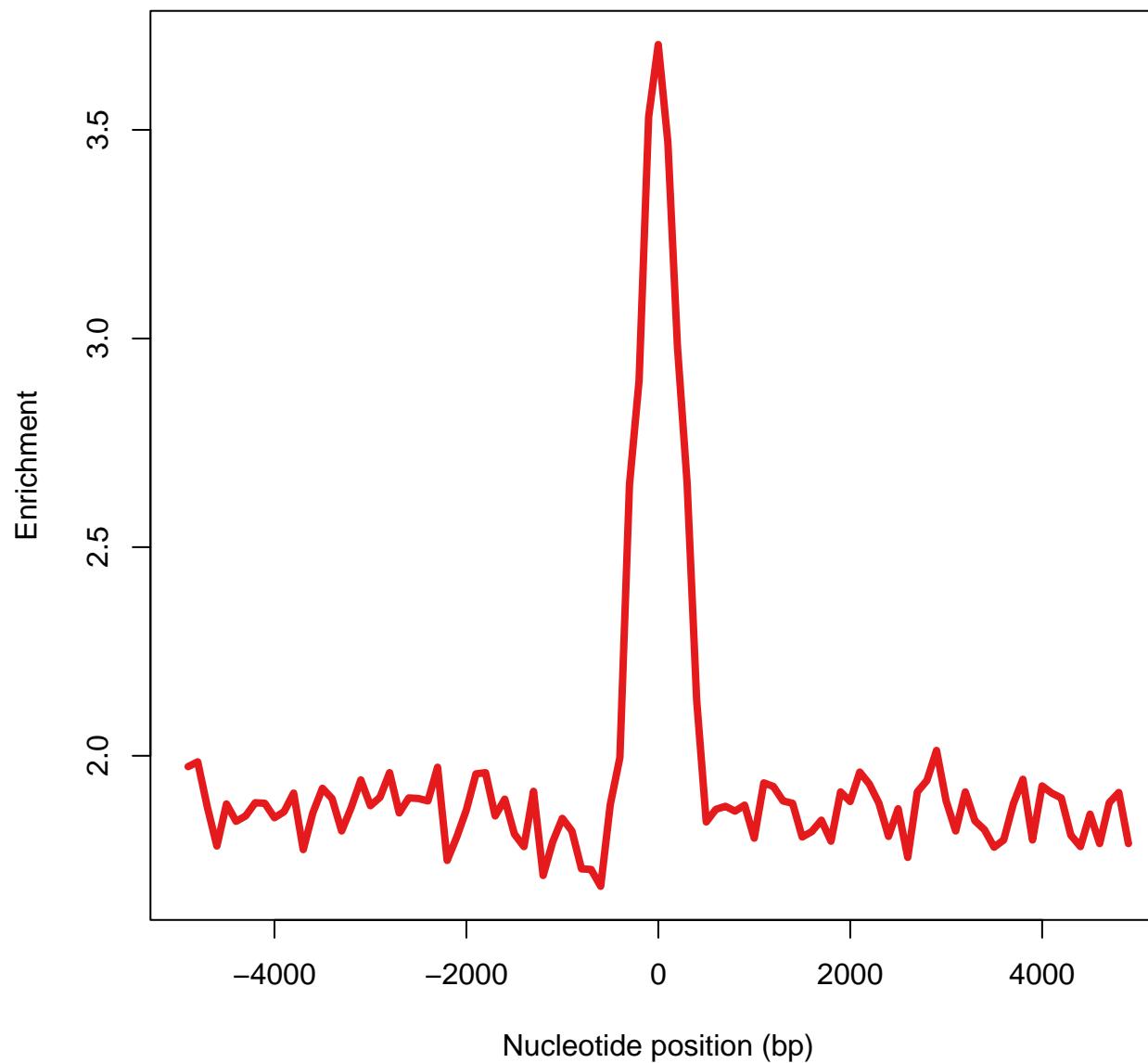
h_2 median distance to genes



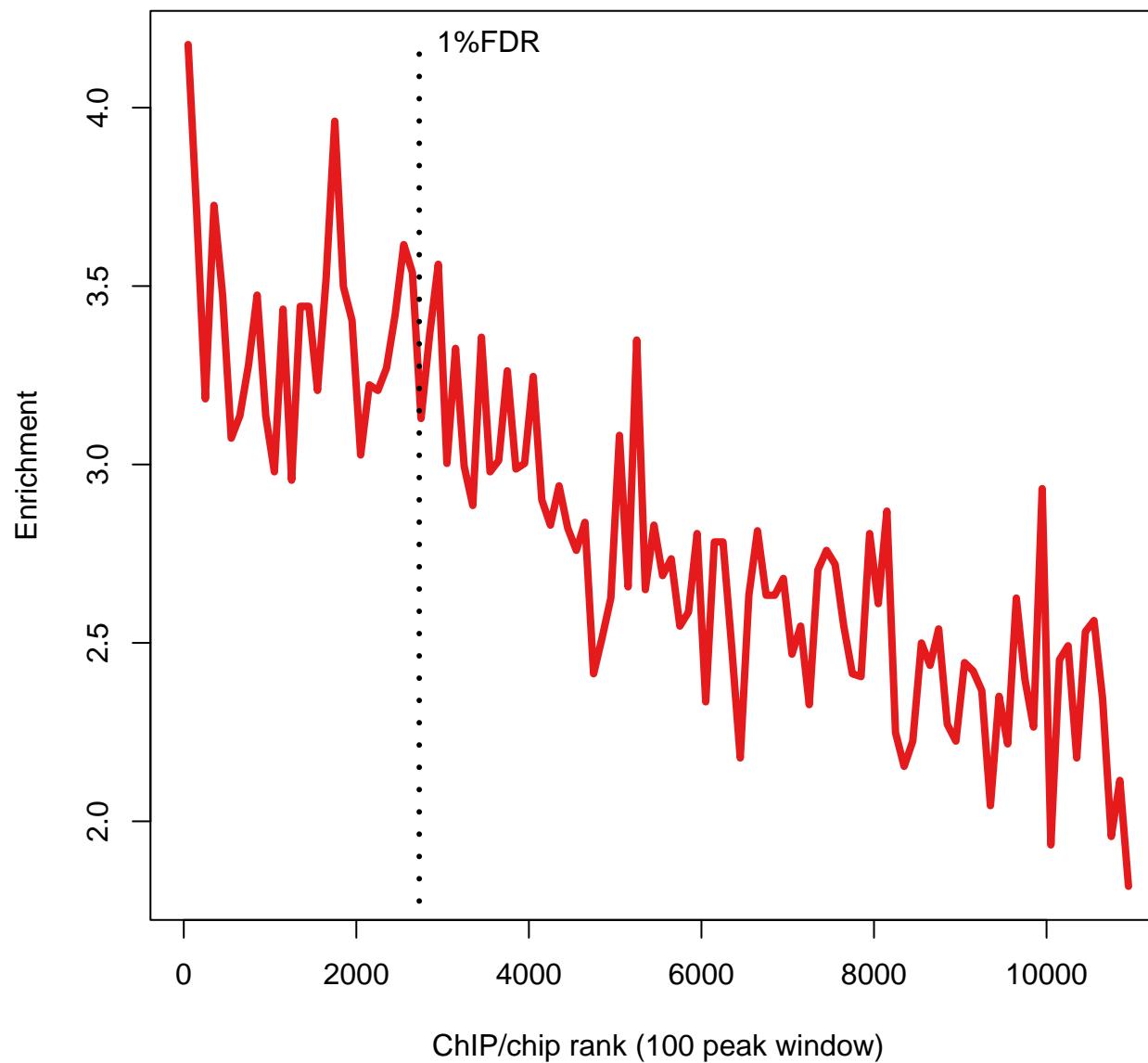
h_2 distribution of number of sites



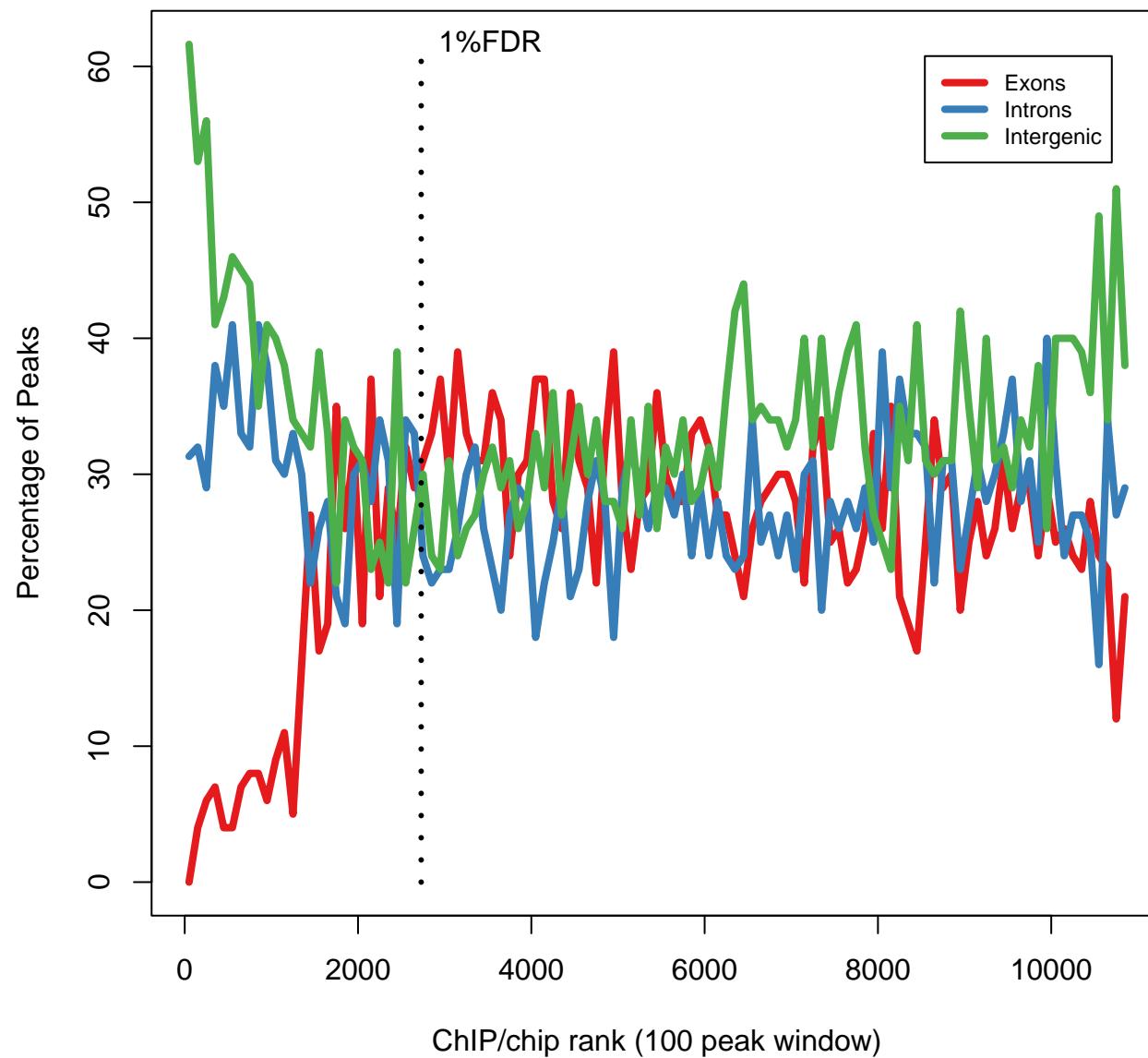
h_2 PWM enrichment across peaks



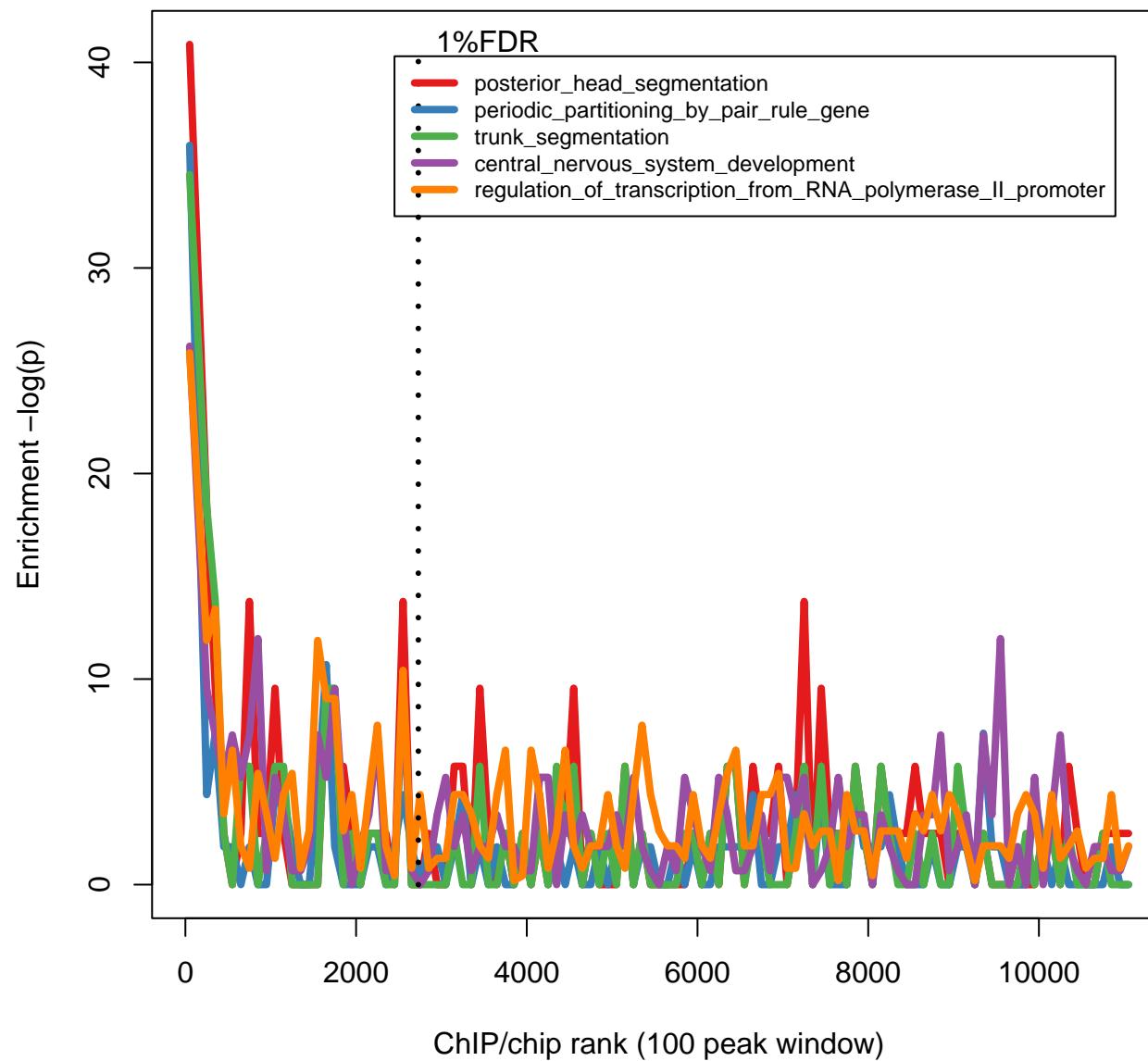
h_2 PWM enrichment down ranks

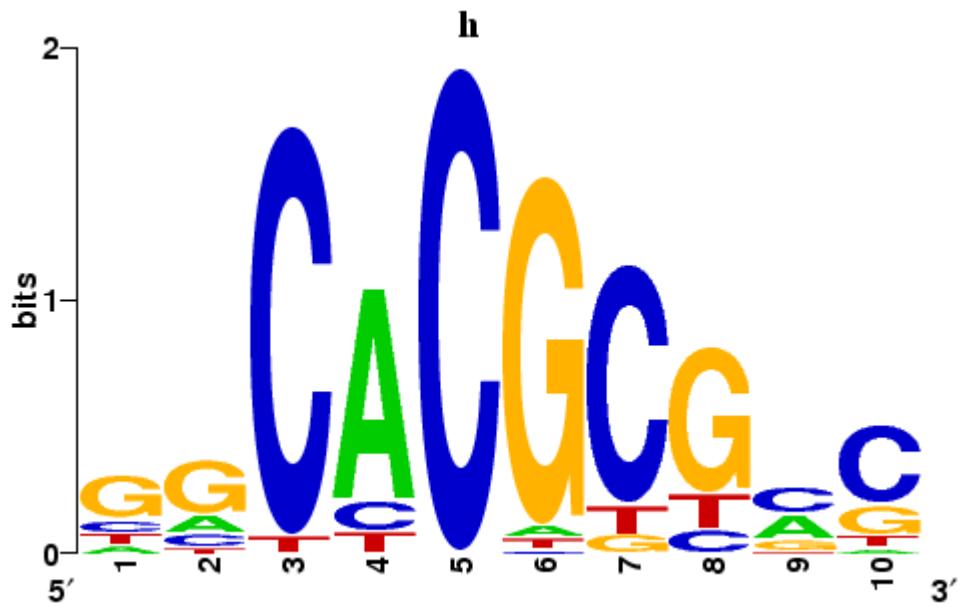


h_2 genomic location

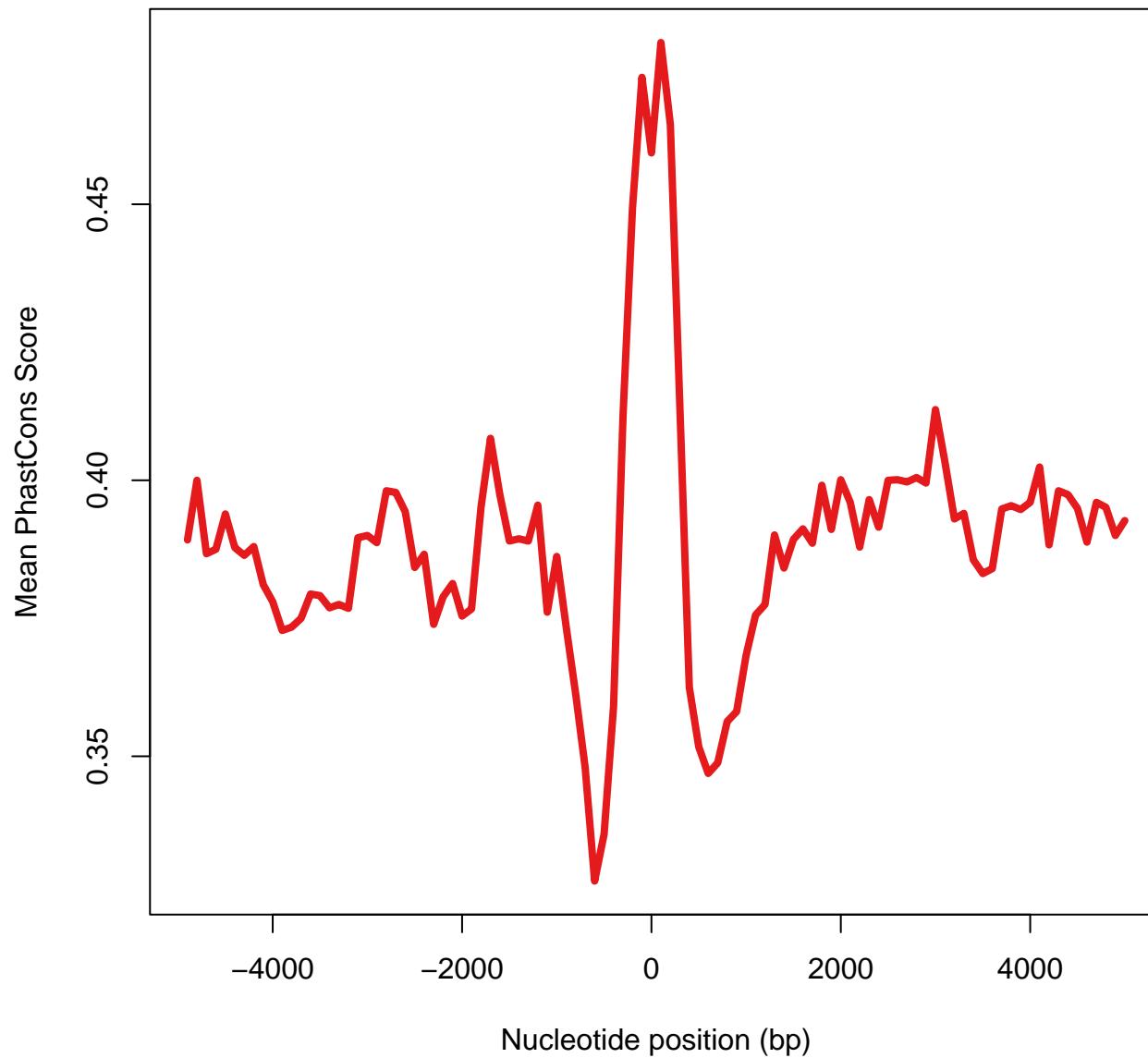


h_2 GO term enrichment (-logP)





h_2 mean PhastCons across peaks



h_2 mean PhastCons down ranks

